

Figure 1

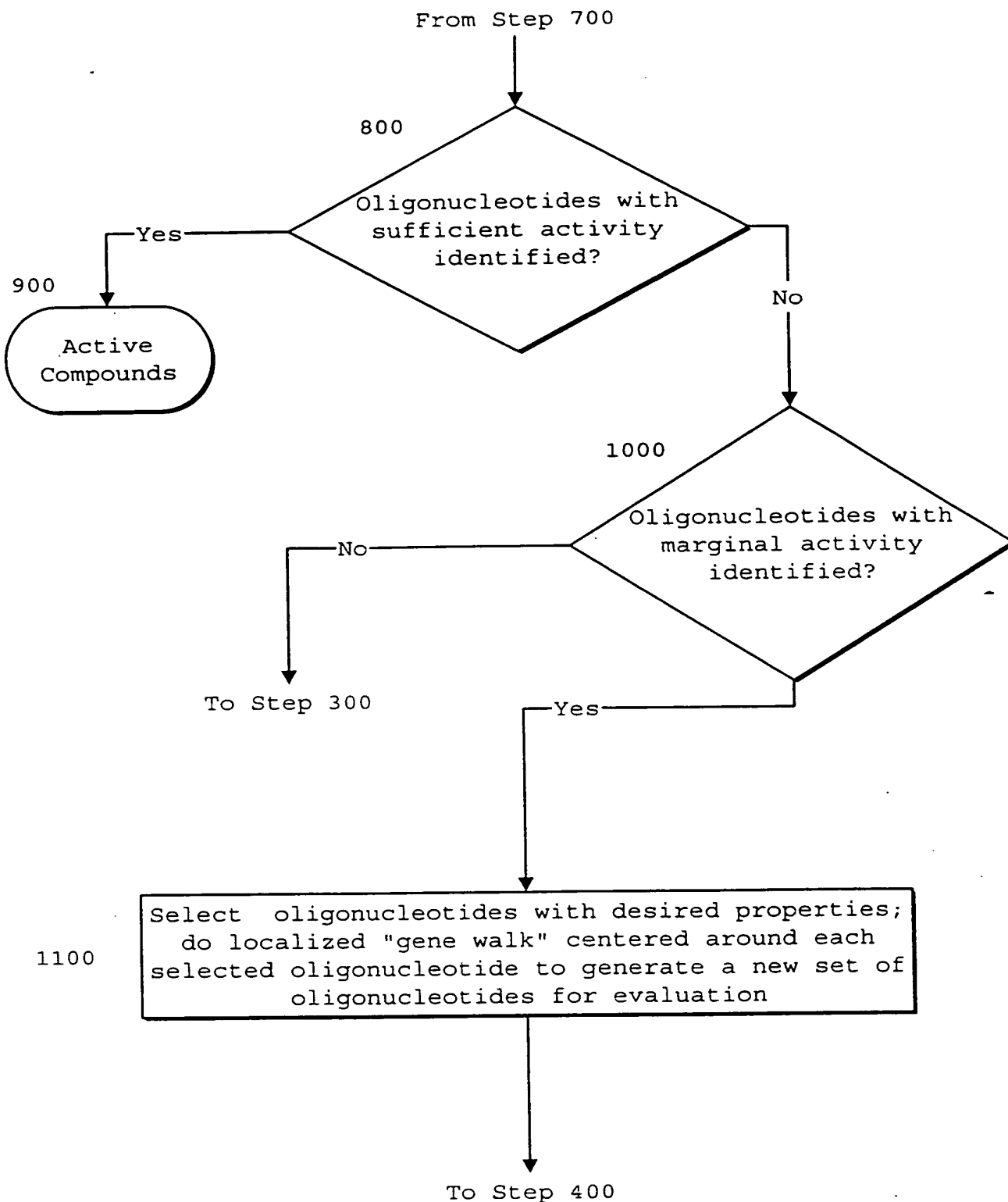


Figure 2

09067638 042898
868240" 8E9/9060

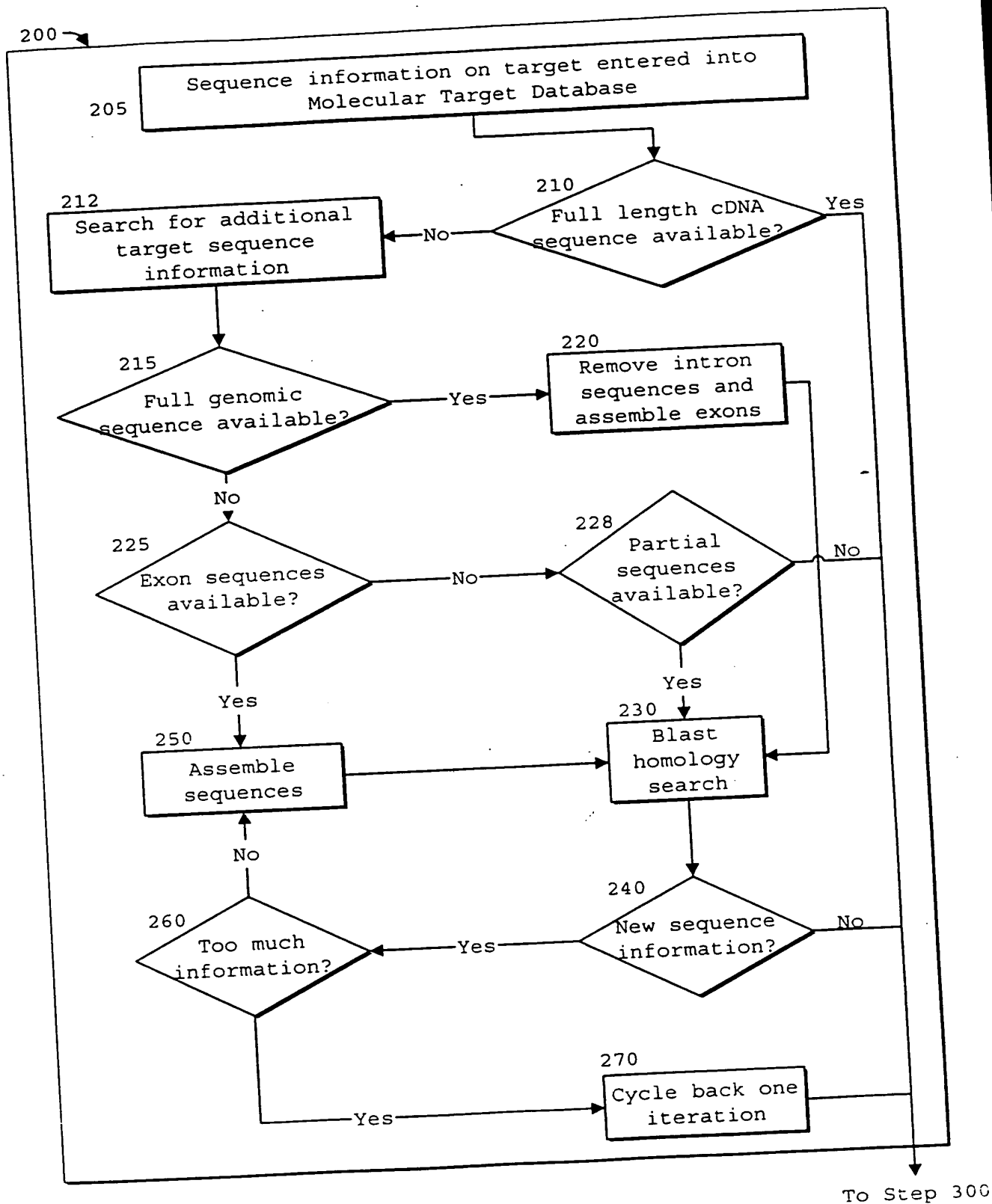


Figure 3

300

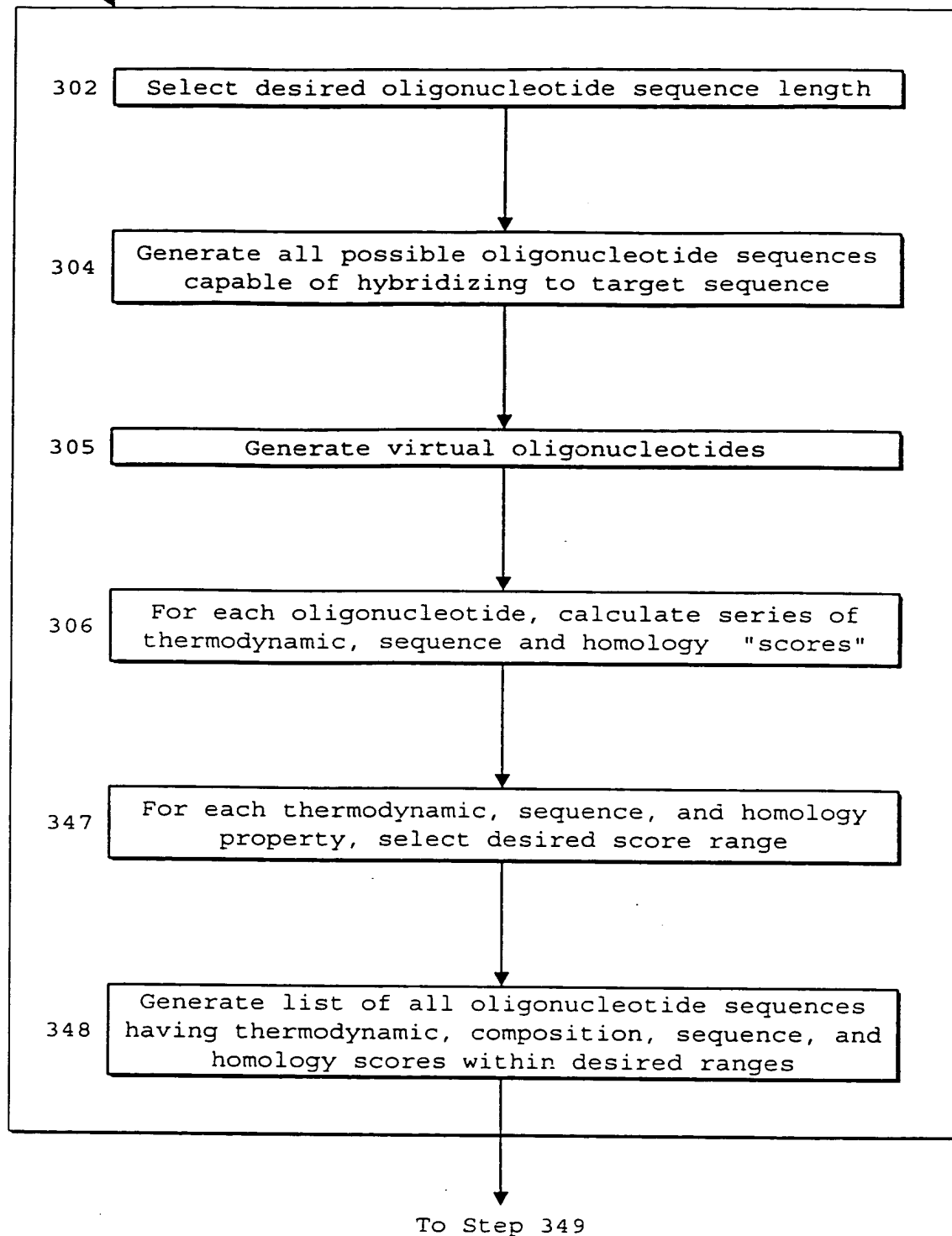


Figure 4

09067638-042898

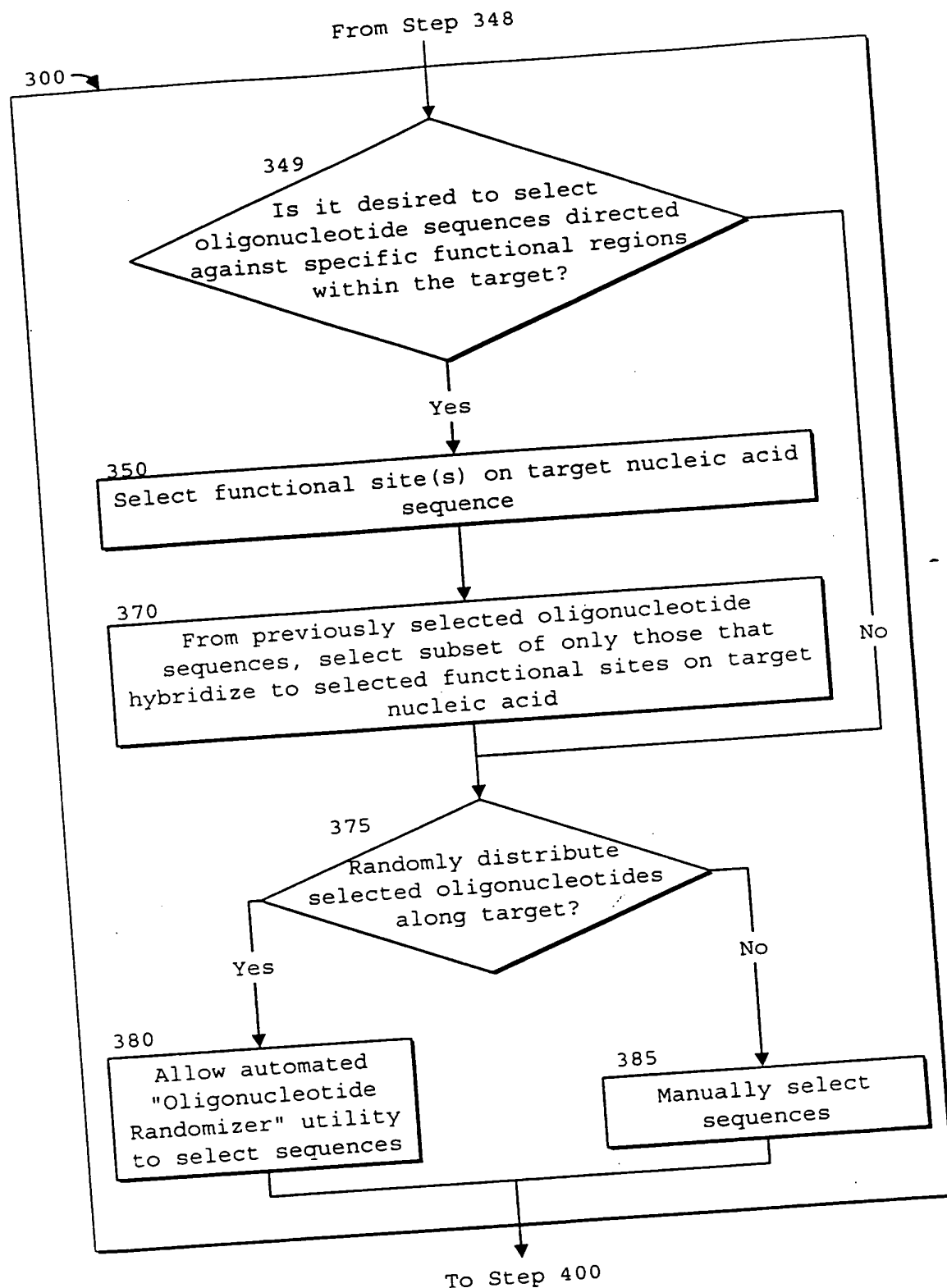


Figure 5

09067638-042898

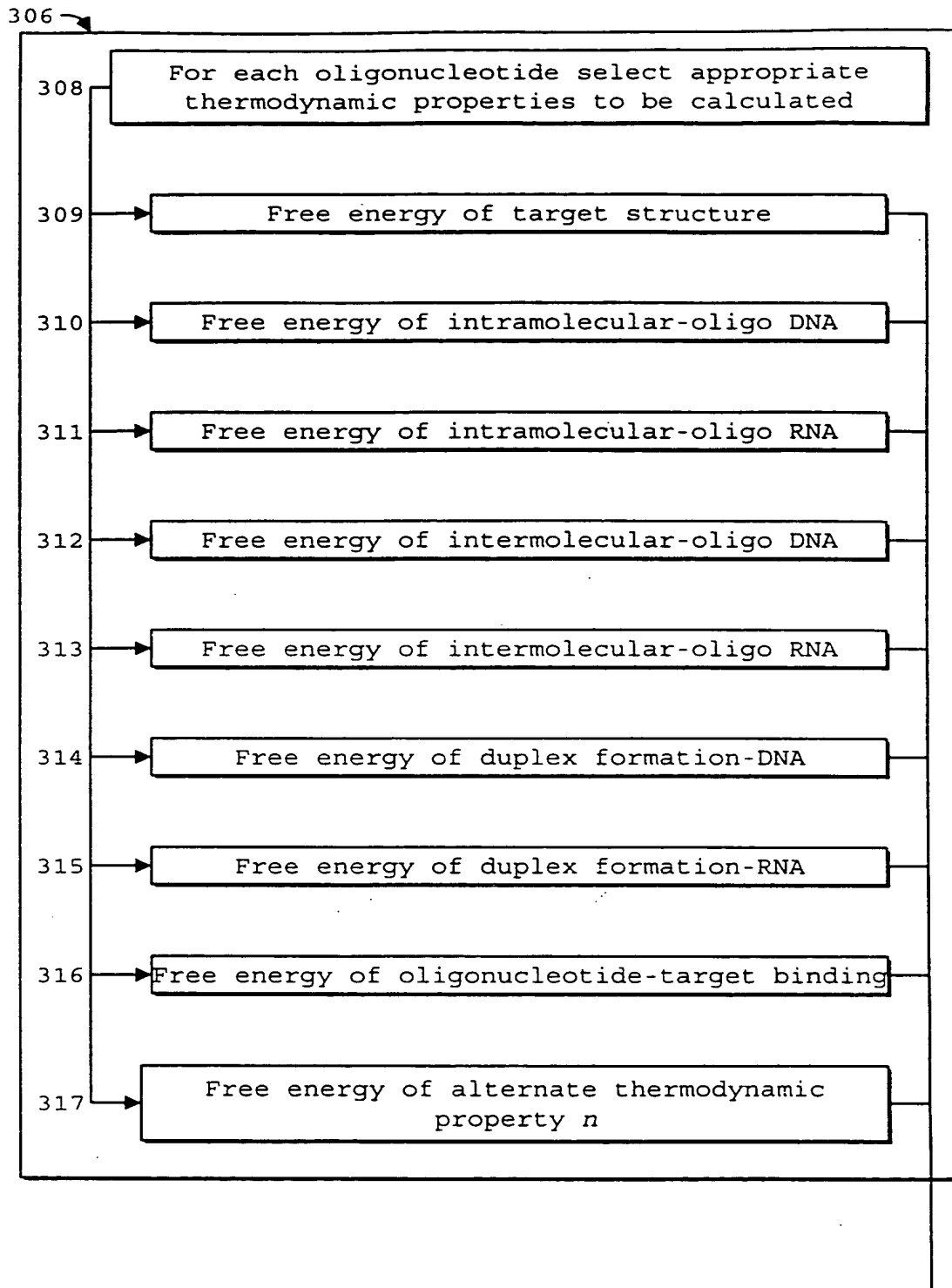
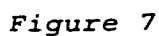
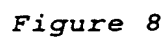


Figure 6

0067538-042893





353	Transcription start site (5' cap)
354	5' UTR
355	Start codon
356	Coding region
357	Stop codon
358	3' UTR
359	5' splice site
360	3' splice site
361	Exon N (N = number of exon)
362	Intron N (N = number of intron)
363	mRNA stabilization signal
364	mRNA destabilization signal
365	Poly-adenylation signal
366	Poly-A addition site
367	Poly-A tail
368	Gene sequence 5' of known pre-mRNA
369	Alternate secondary structure property n

Figure 9

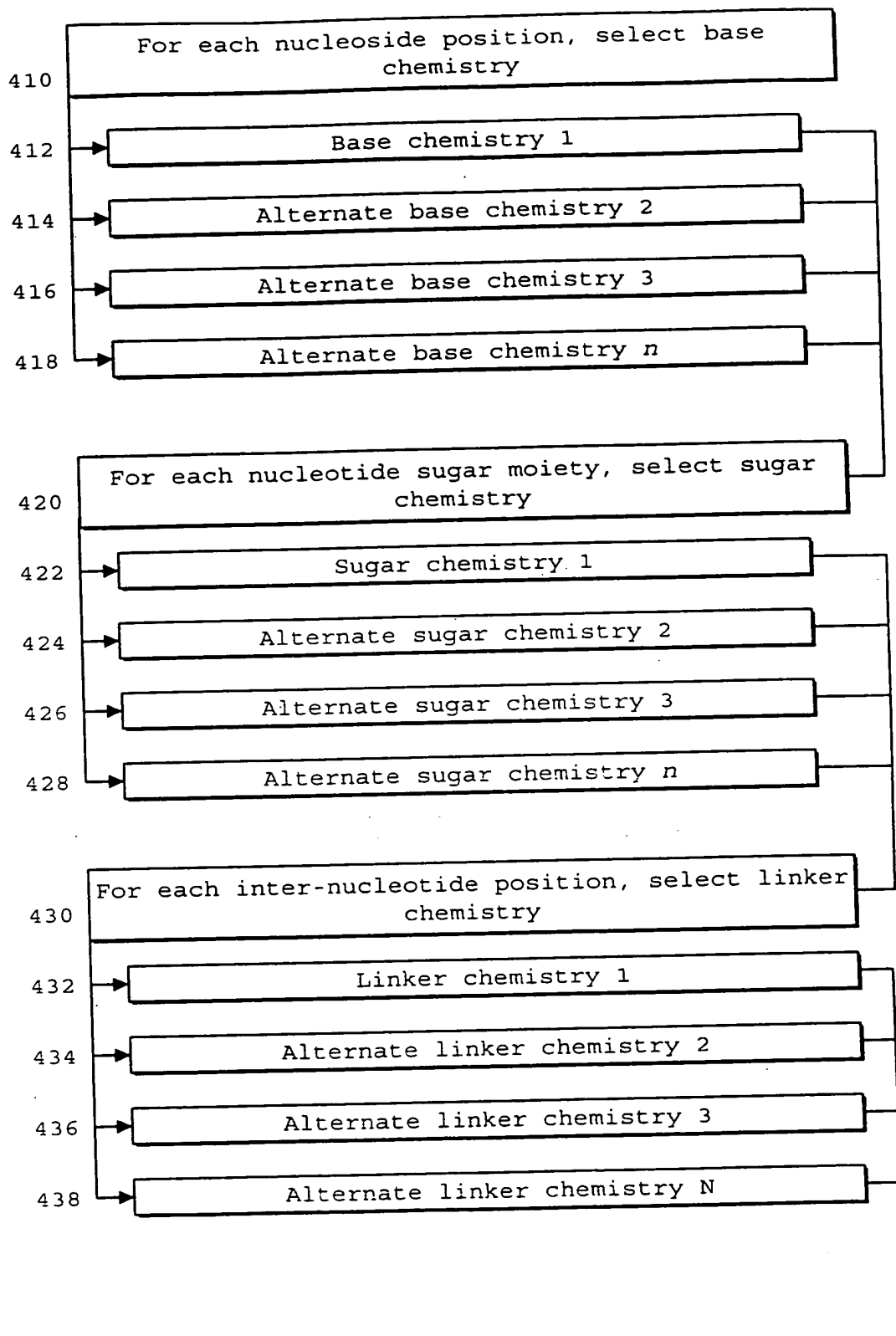


Figure 10

To Step 440

00067638 042898

09067638 042890

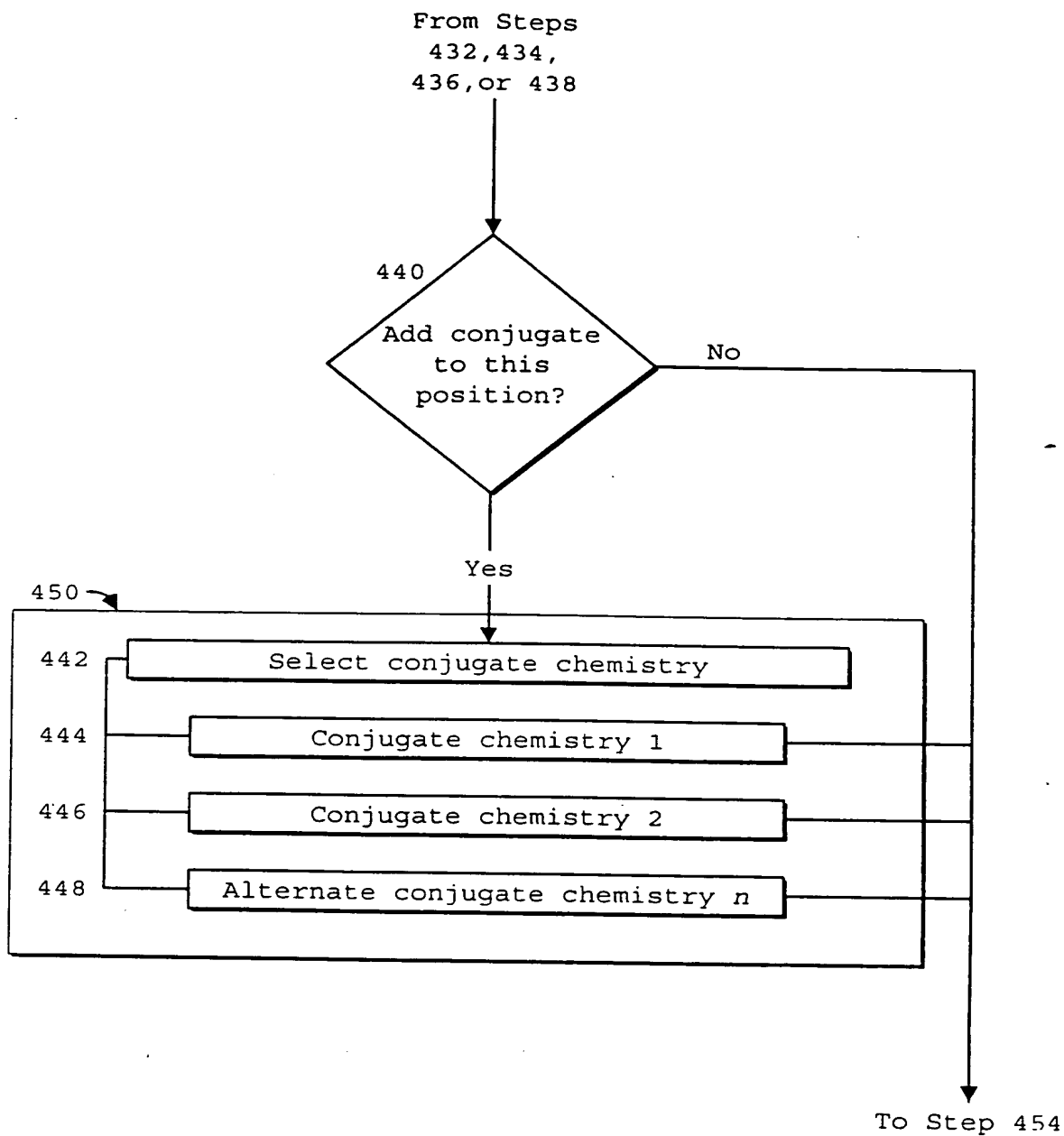


Figure 11

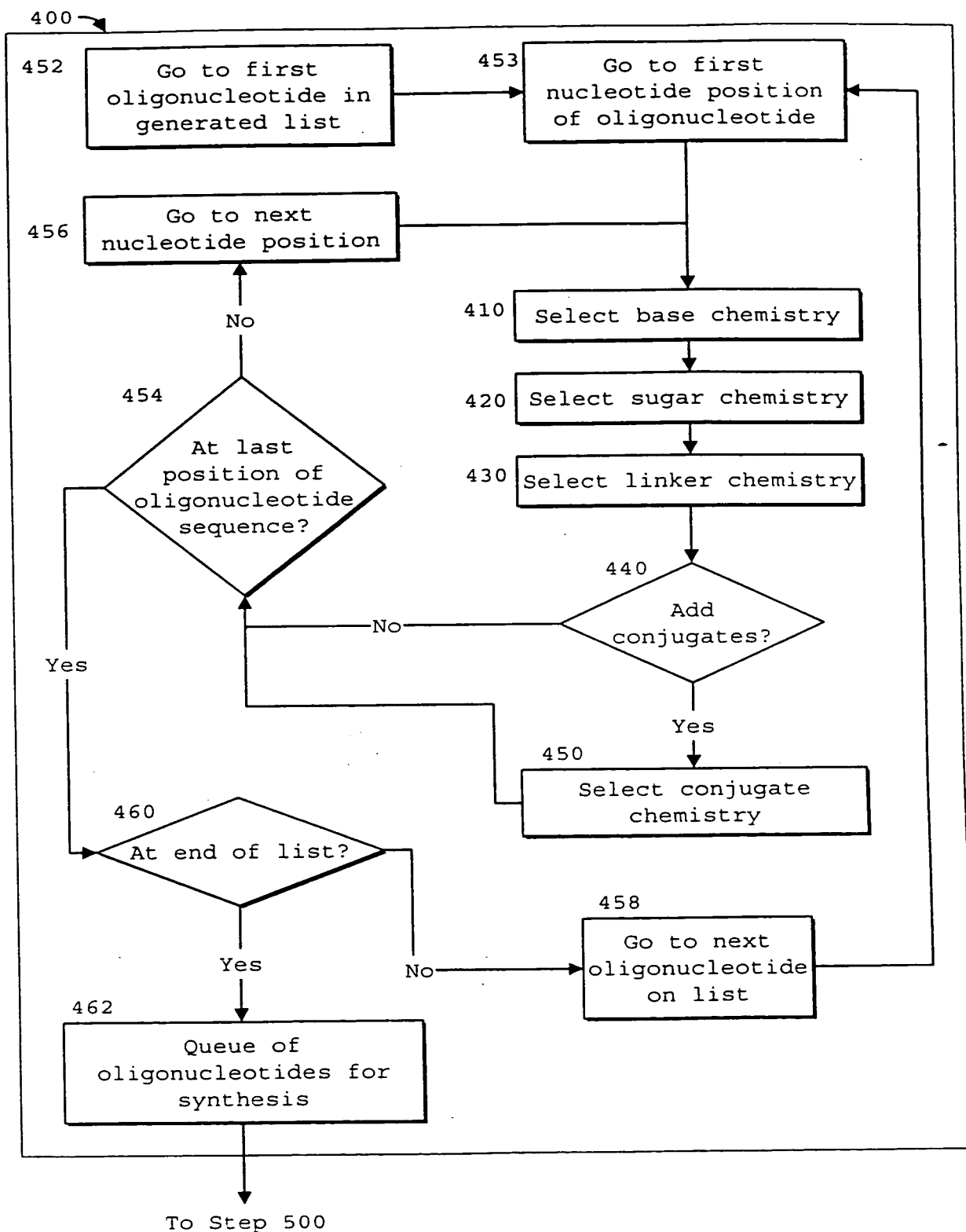


Figure 12

09067639 042898

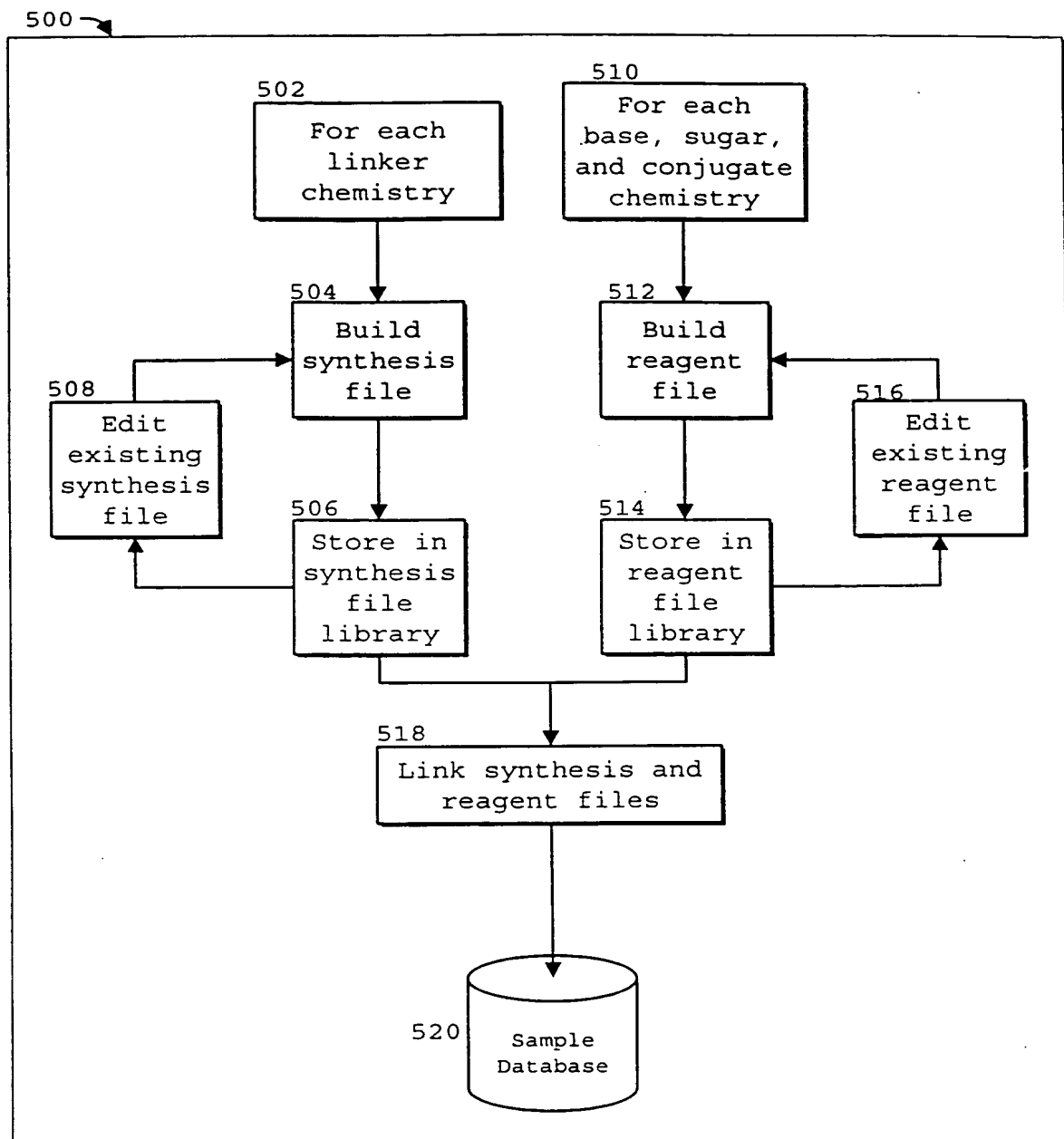


Figure 13

00067638-042890

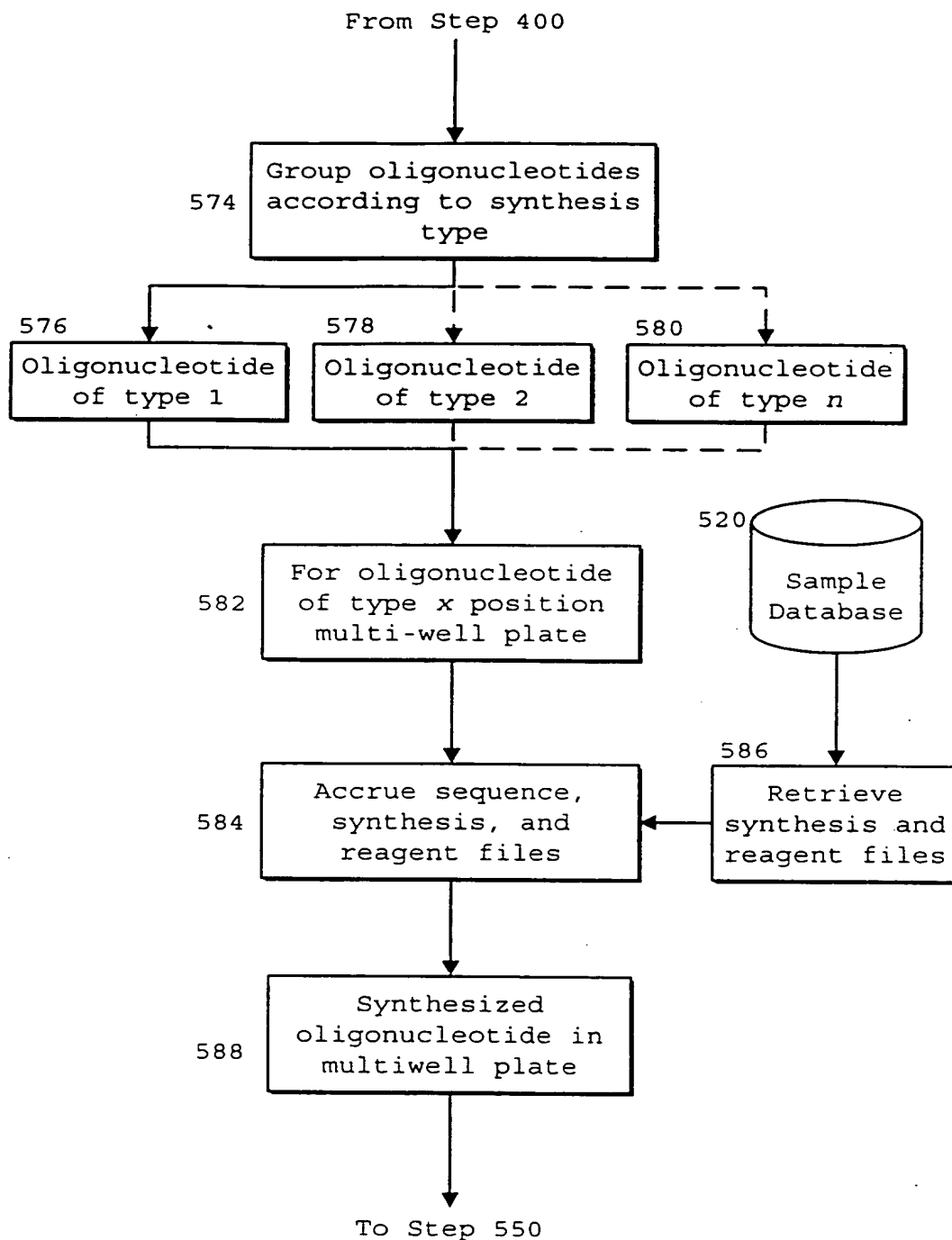


Figure 14

00067638 042898

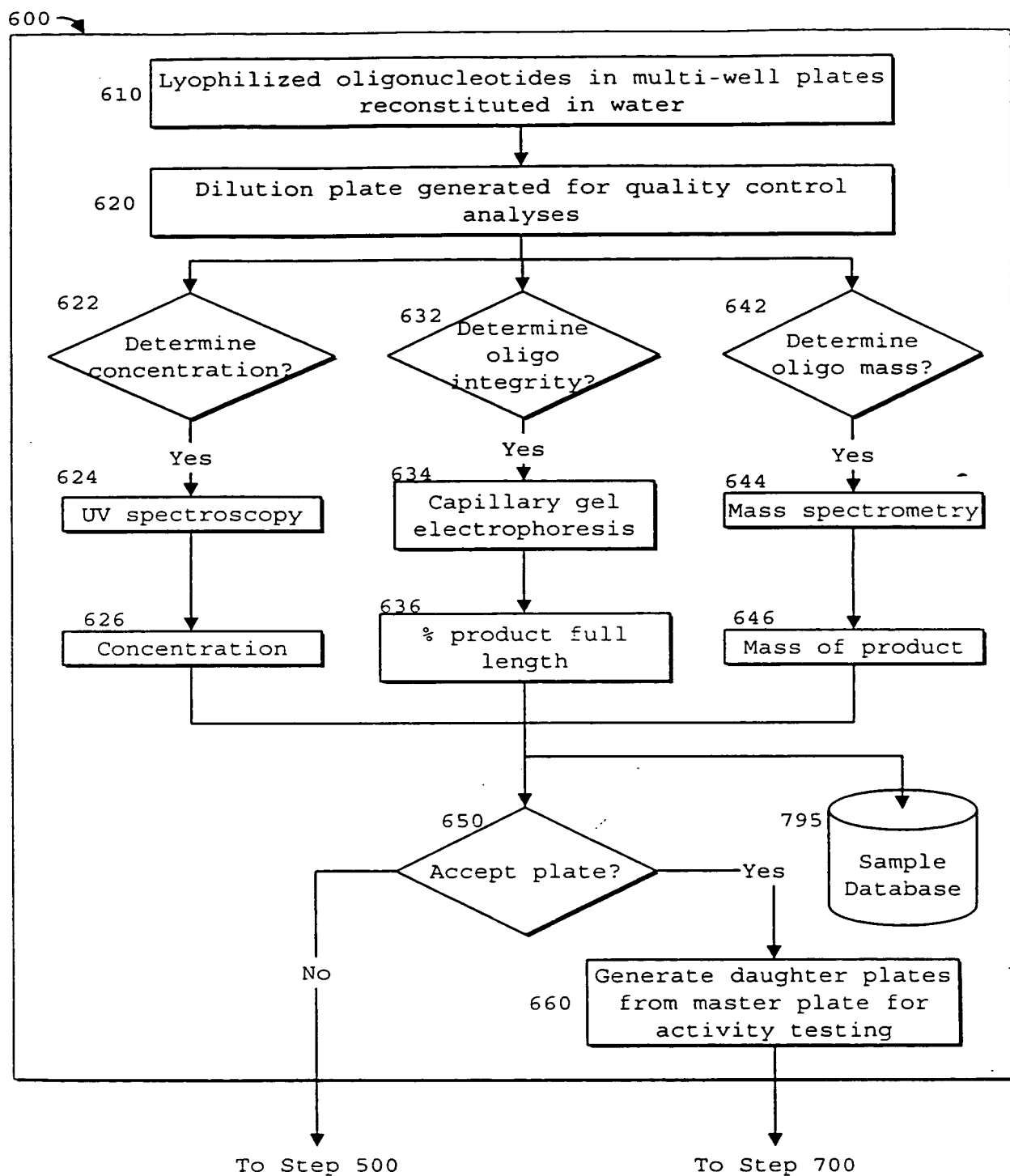


Figure 15

00067638-042898

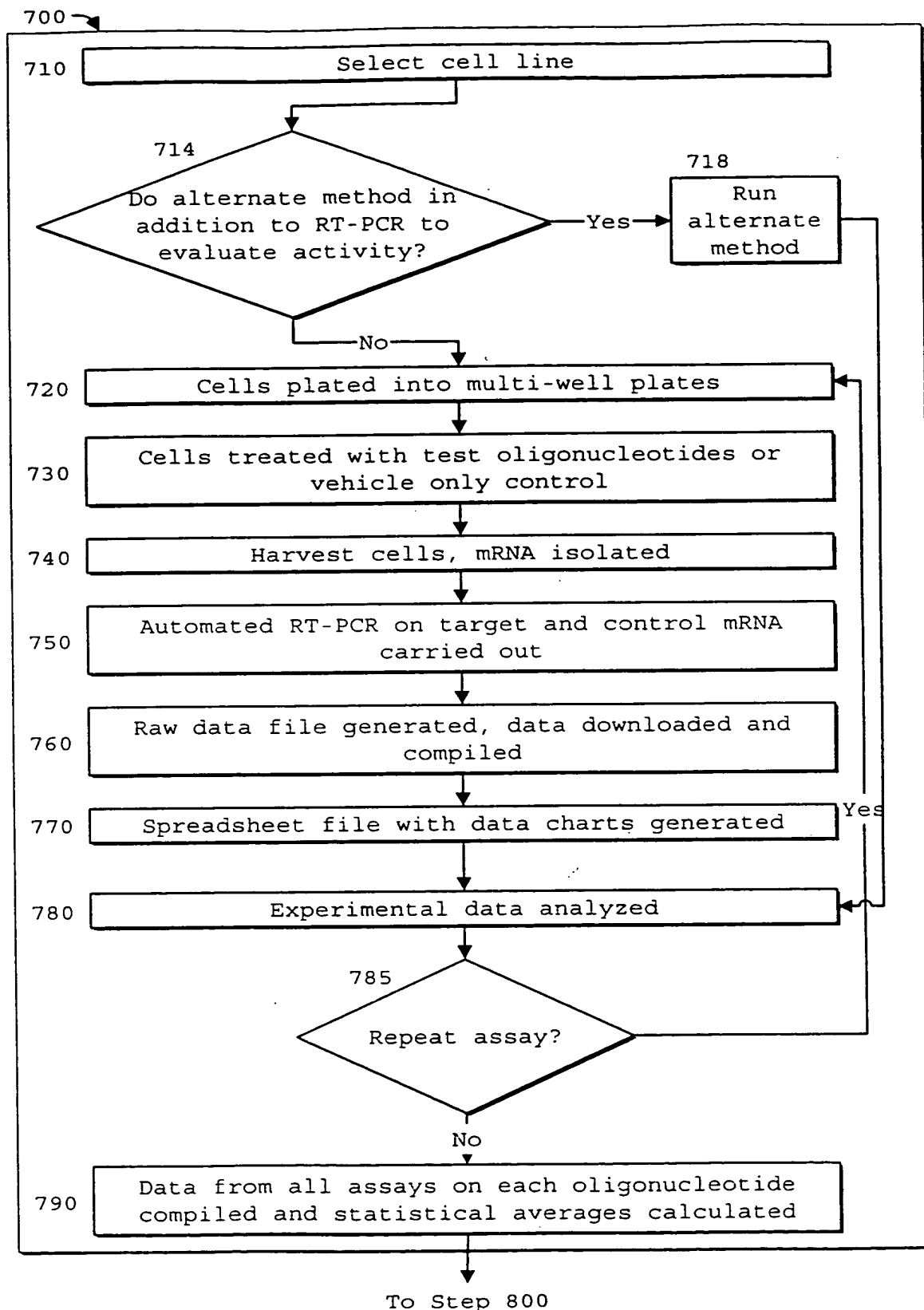


Figure 16

0067638-042899

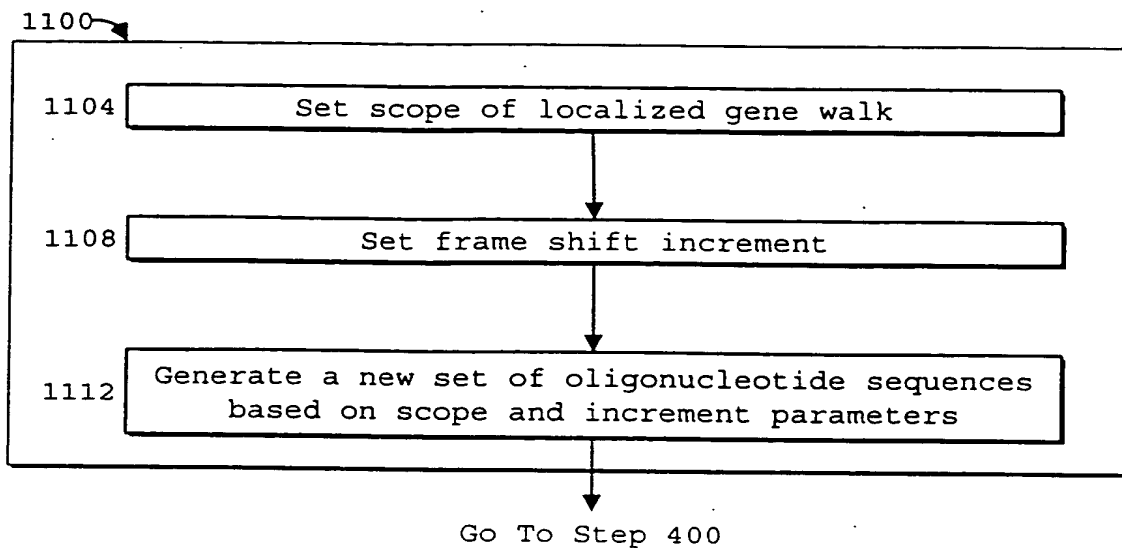


Figure 17

006240-88929050

868240" 8E923060

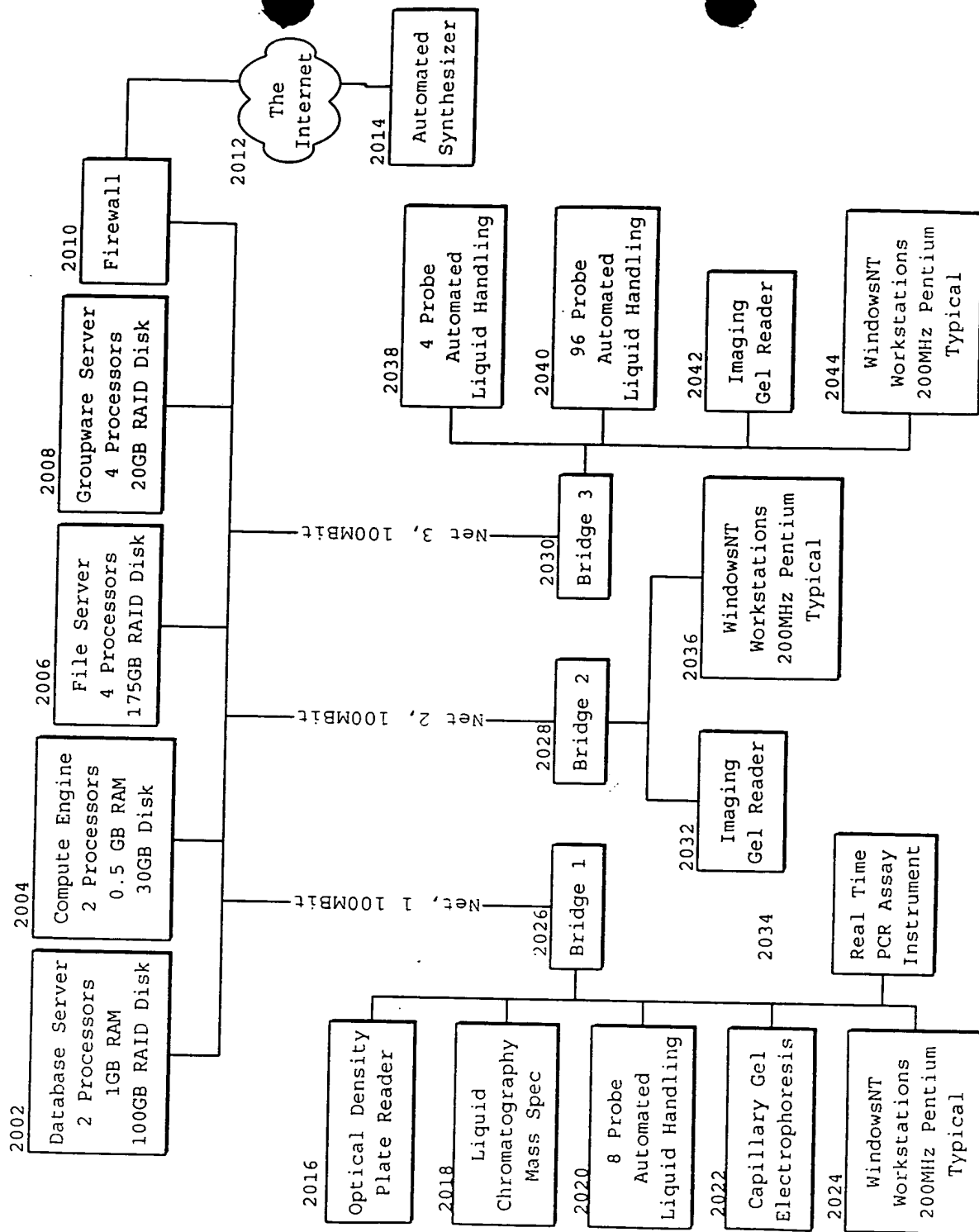


Figure 18

Figure 19

36340" 36929060

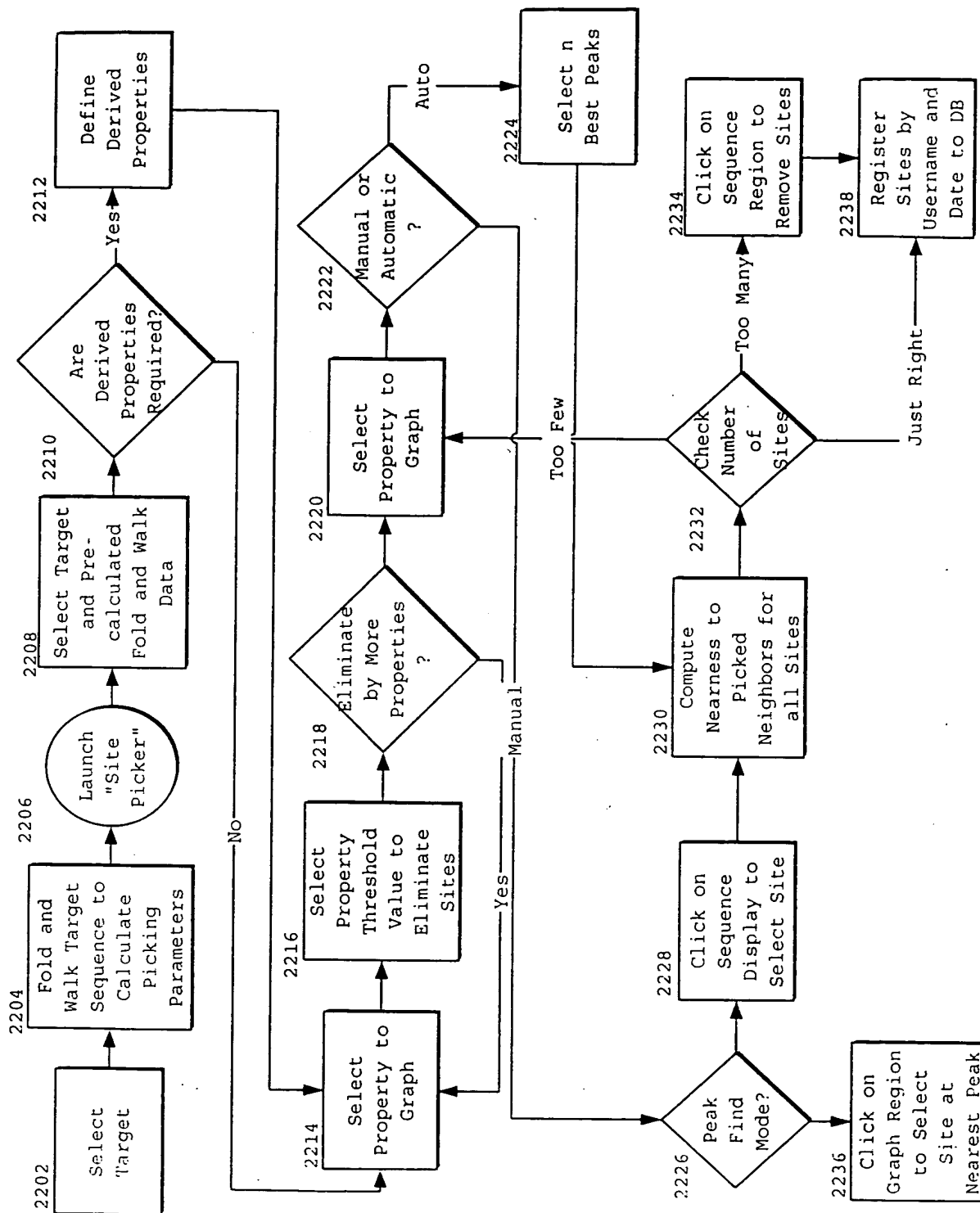
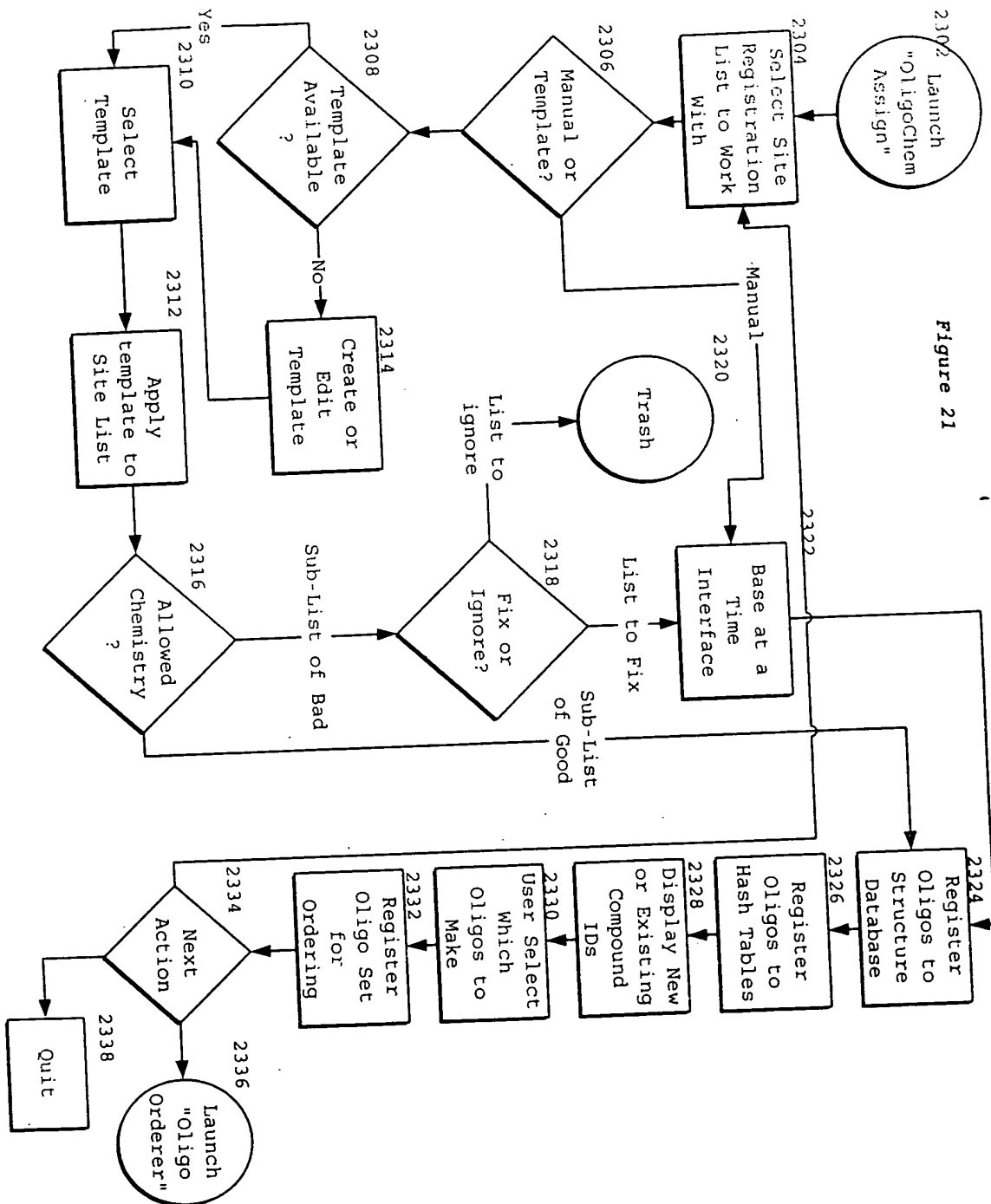


Figure 20

Figure 21



09067638 . 042398

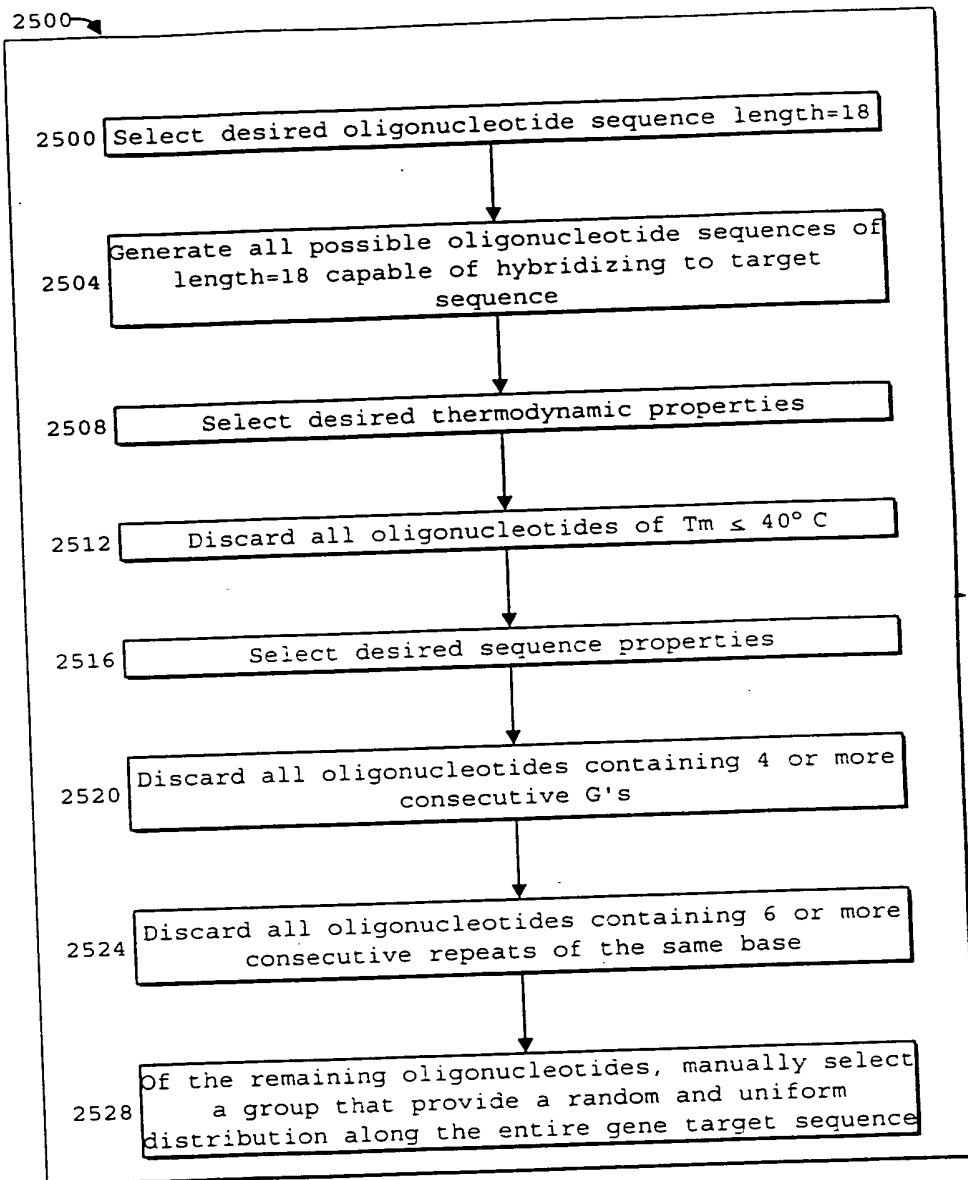


Figure 22

09067638-042888

868240" 86929060

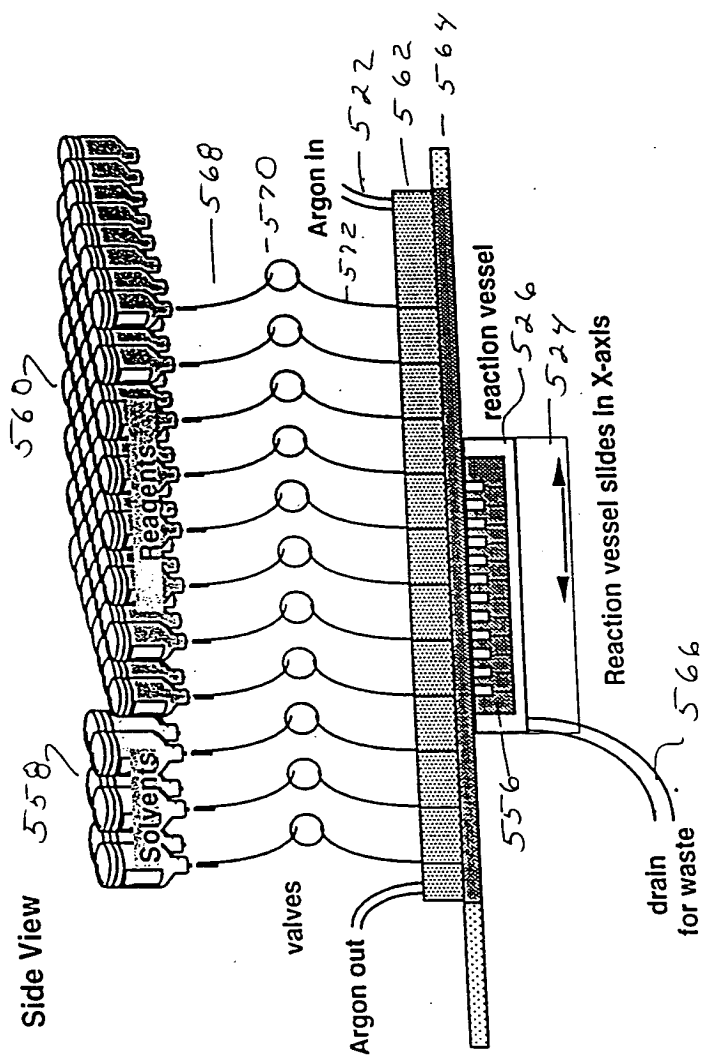


Figure 23

1306240" 8E929060

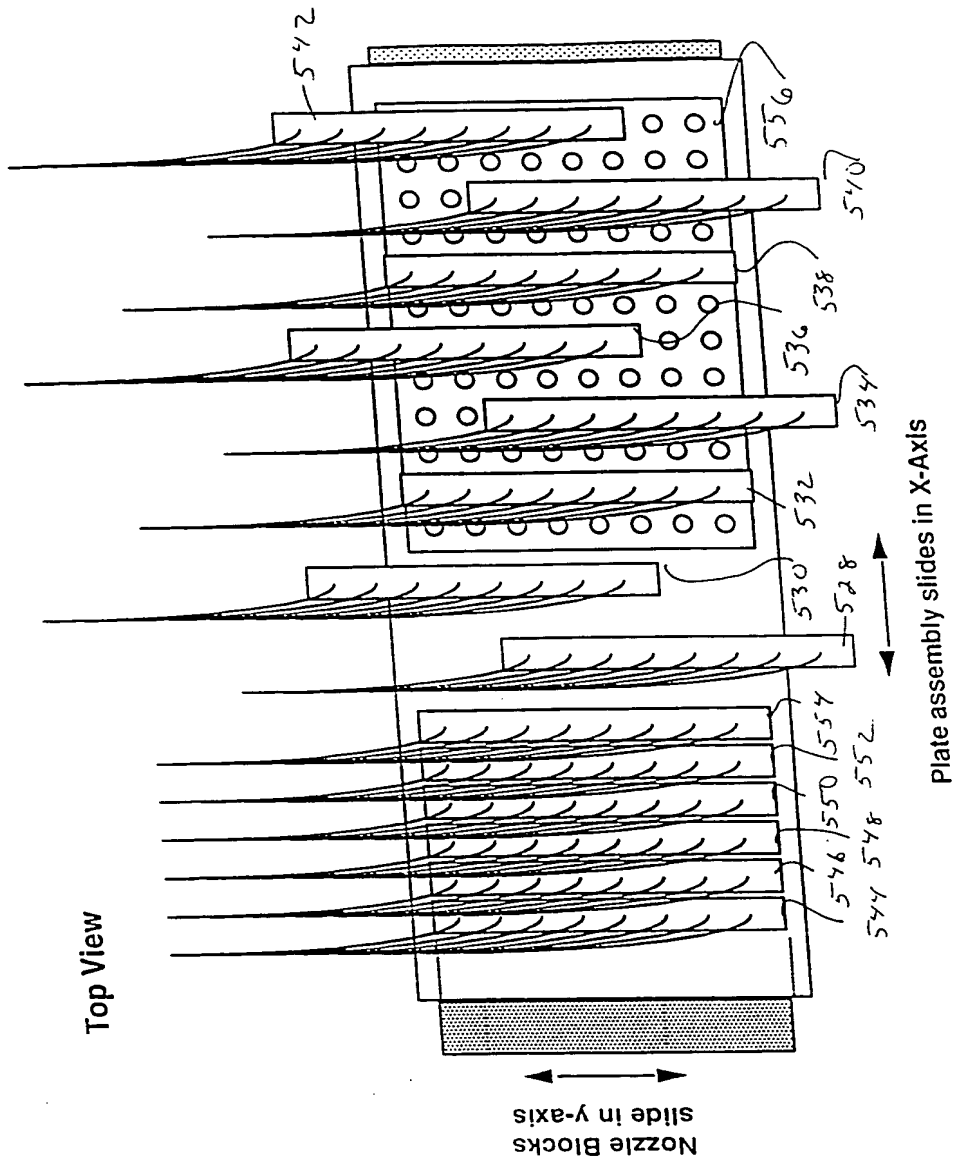


Figure 24